

1/18

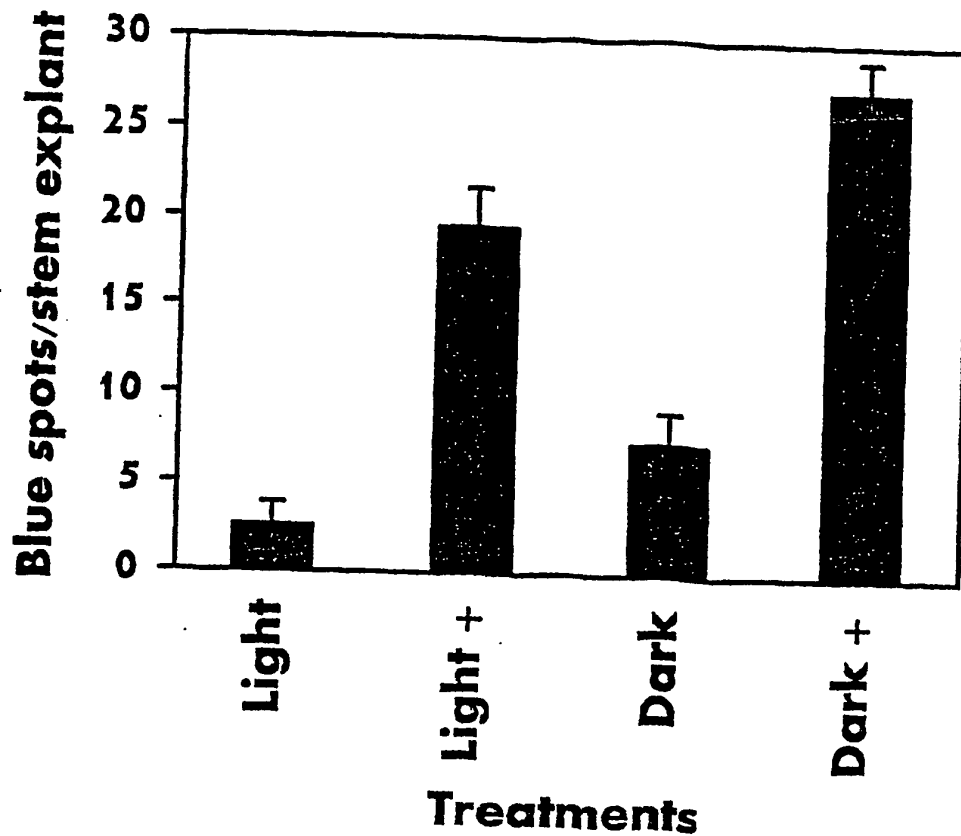


Fig. 1

2/18

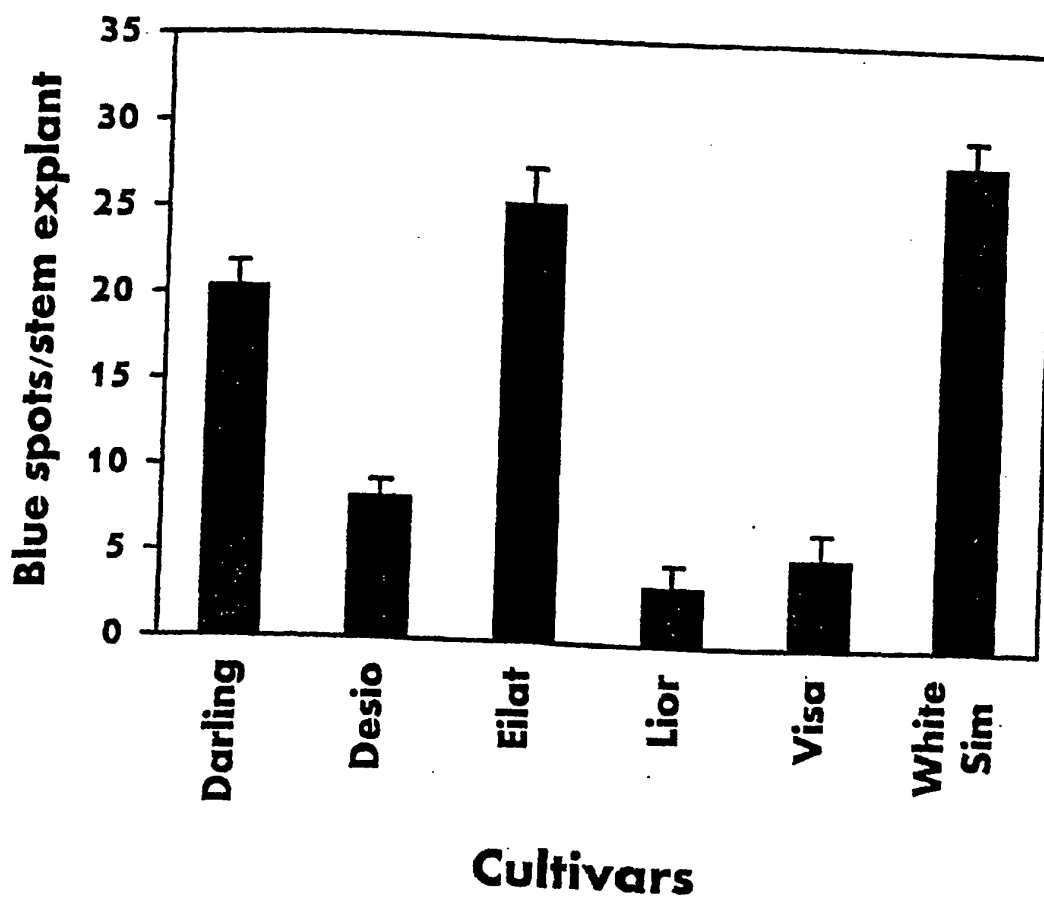


Fig. 2

3/18

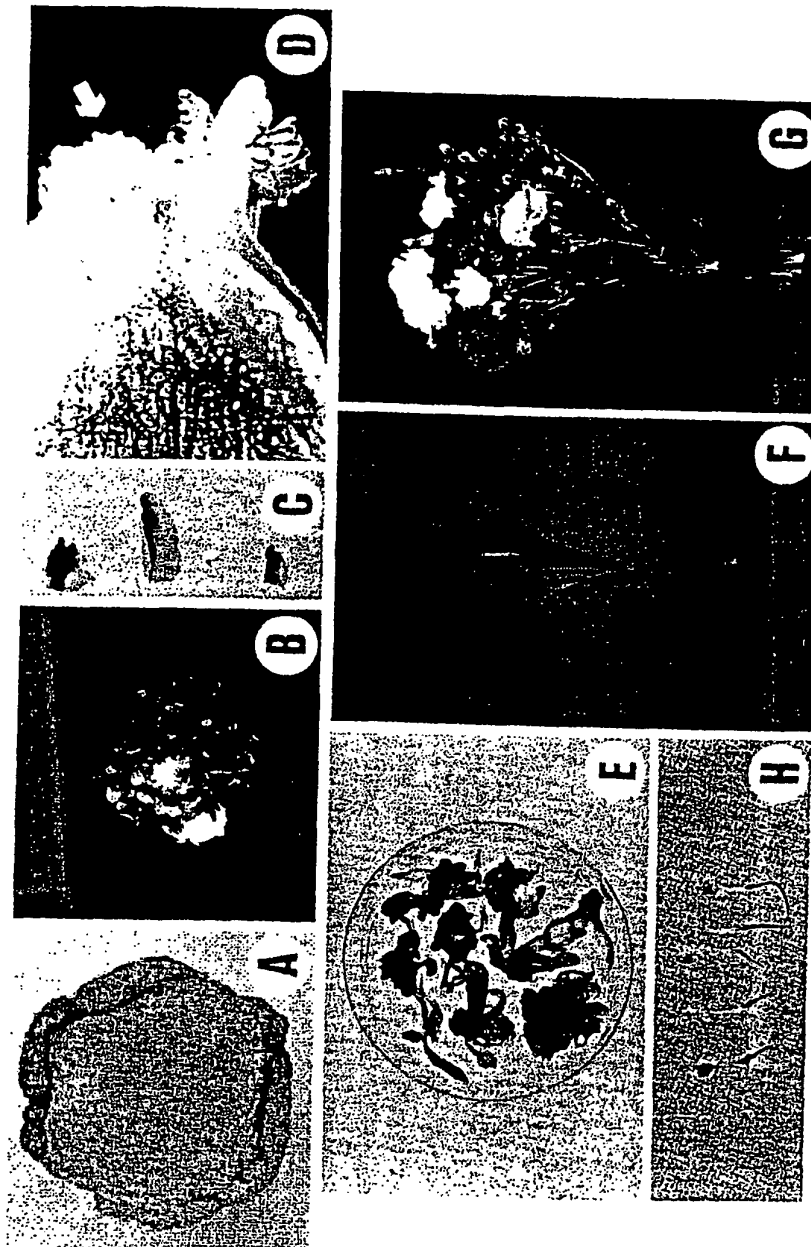


Fig. 3

206270" 94TH T660

4/18

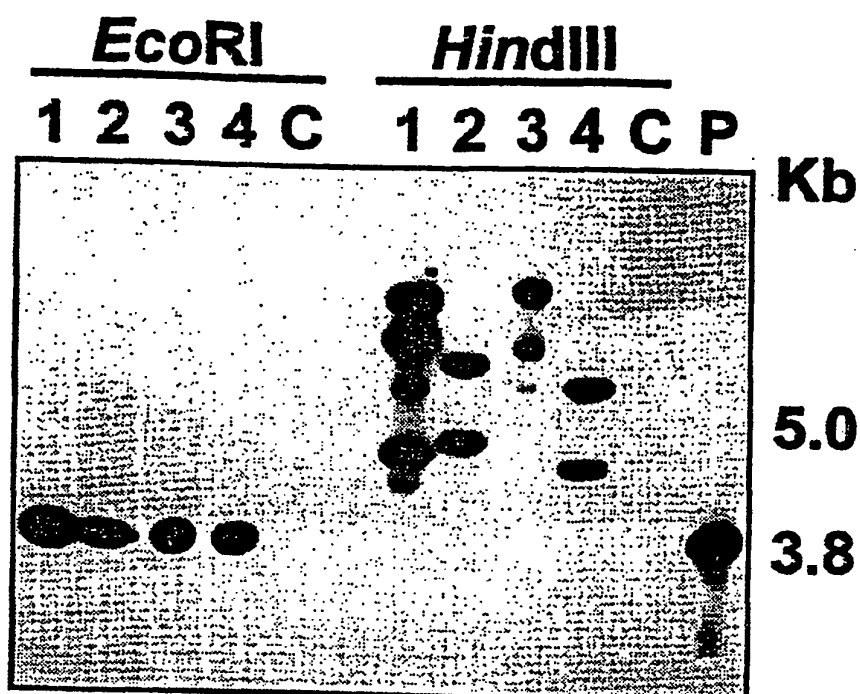


Fig. 4

206270" 94747650

5/18

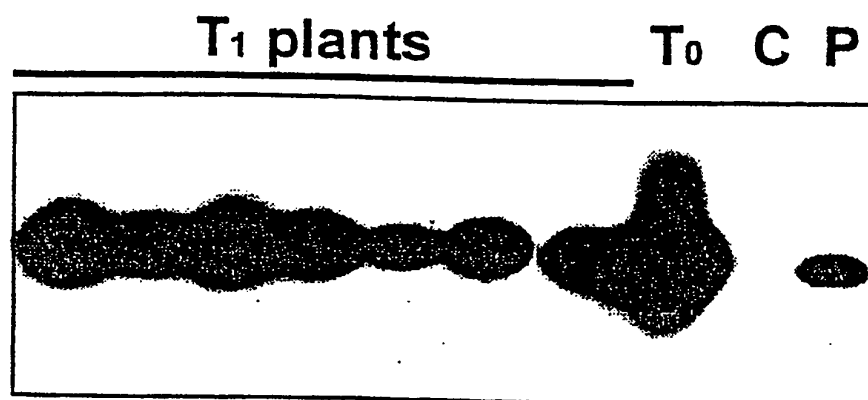


Fig. 5

800bp

09/914146-012900
2062T0" 94T+T660

6/18

206210" 9444660

1 ATGGCTGAAG ACGACCTGTG TTCTCTCTTT TTCAAGCTCA AAGTGGAGGA TGTGACAAGC
61 ACGGATGAGC TAGCTAGACA CATGAAGAAC GCCTCAAATG AGCGTAAACC CTTGATCGAG
121 CCGGGTGAGA ATCAATCGAT GGATATTGAC GAAGAAGGAG GGTGGGTGG CCACGGGCTG
181 CTGTACCTCT ACGTCGACTG CCCGACGATG ATGCTCTGCT TCTATGGAGG GTCCCTGCCT
241 TACAATTGGA TGAAGGGCG ACTCCTCACC AACCTTCCCC CGTACCAGCA TGATGTGACT
301 CTCGATGAGG TCAATAGAGG GCTCAGGCAA GCATCAGGTT TTTTCGGTTA CGCGGATCCT
361 ATGCGGAGCG CCTACTTCGC TGCATTTTCT TTCCCTGGGC GTGTCAATCA GCTGAATGAG
421 CAGATGGAGC TAACTTCGAC AAAGGGAAG TGCTGACAT TCGACCTCTA TGCCAGCACC
481 CAGCTTAGGT TCGAACCTGG TGAGTTGGTG AGGCATGGCG AGTGCAAGTT TGCAATCGGC
541 TAATGGTTAG TCGATGGGCT GACGAGTTTG ATGTCAGGAG AAGCTGAGTG TGTCACCTGT
601 TTCCCTTTAA GAAGTATTAA TGTAATAAAA ATCAAGATCT GGTTTAATA CTGGATACTT
661 GATTTCATCG CGCTTTTTTT GAATAAATGT TTGTTGTCTT GACTTTAAGA TATCCTTTGA
721 AATTGCGGTT ATTCGTATTT CGCTTTTGGT TATTTCCAAA AGACTTTGCT CAGTAAGATC
781 AAACGTTTGT ATTTCICCGG GCCACAATAT TTGACCTATA TGCCTGGCC CACGCGCCGC
841 AATAGATGAA AATTGCCAAA ATTAGCTATC GGTCTTCTGA AAAGAAGGGC CGACATGTTT

Fig. 6

7/18

206210" SHHTE60

901 TCATAGACCA TGCAAAGTCA TACTACCTGA AACTGATAAA TAACGACAAA GAAAGTAGCC
961 TATTAAAAG TCGCTATAGC ATGAATT

Fig. 6 (cont.)

8/18

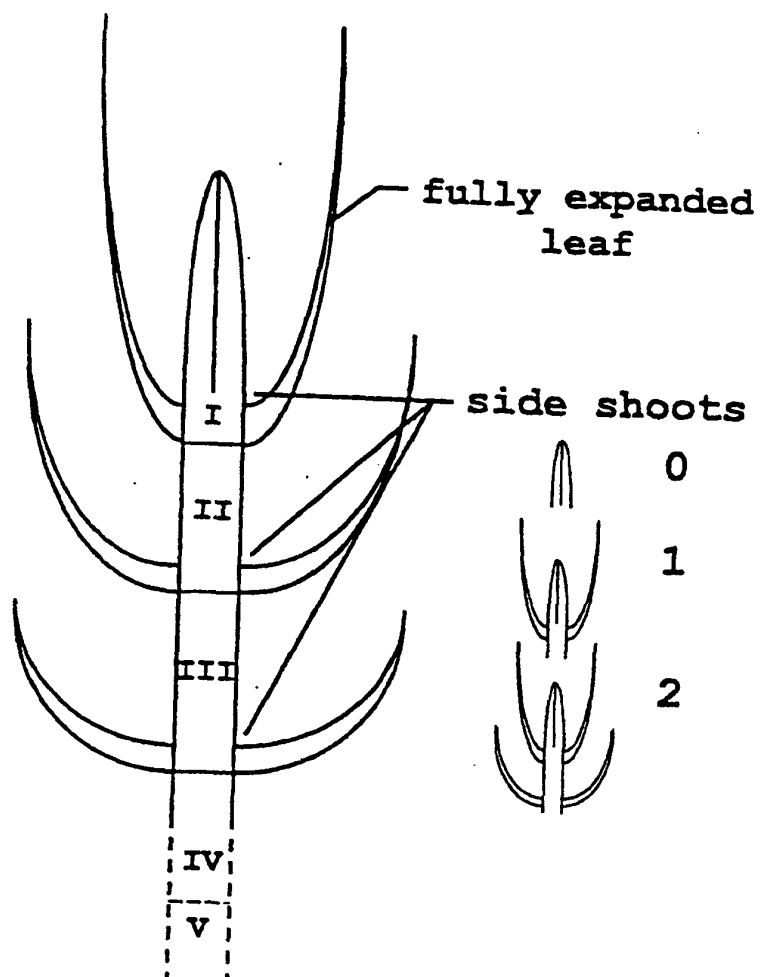


Fig. 7

9/18

206270" 94747660

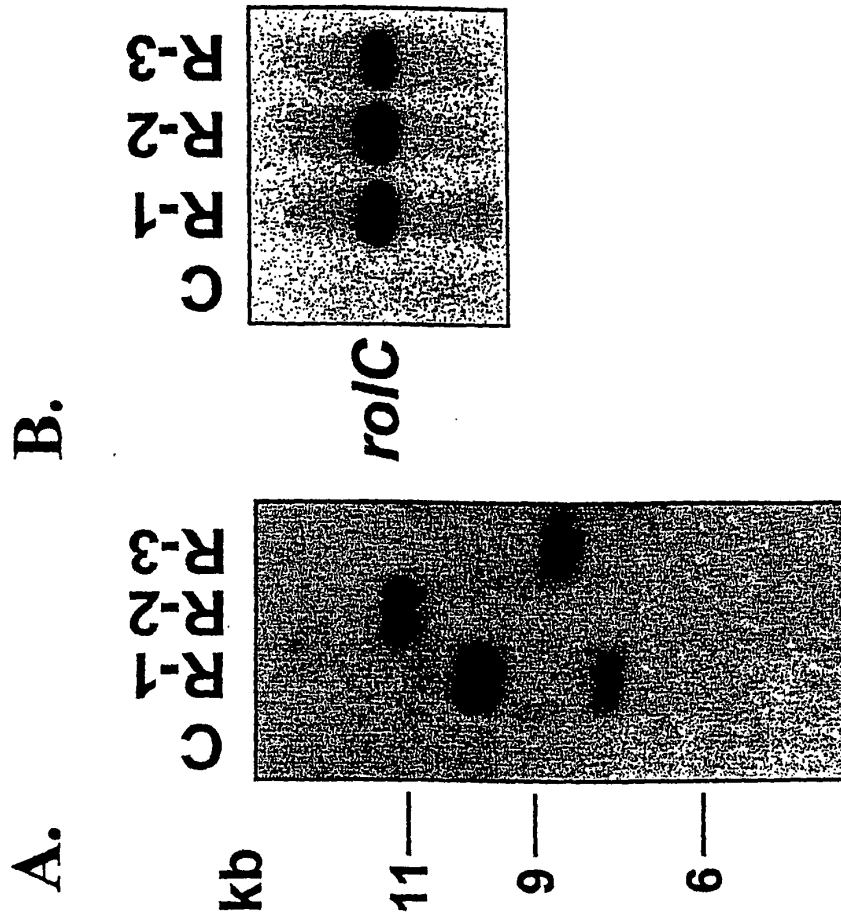


Fig. 8

10/18

206270" 94TH660

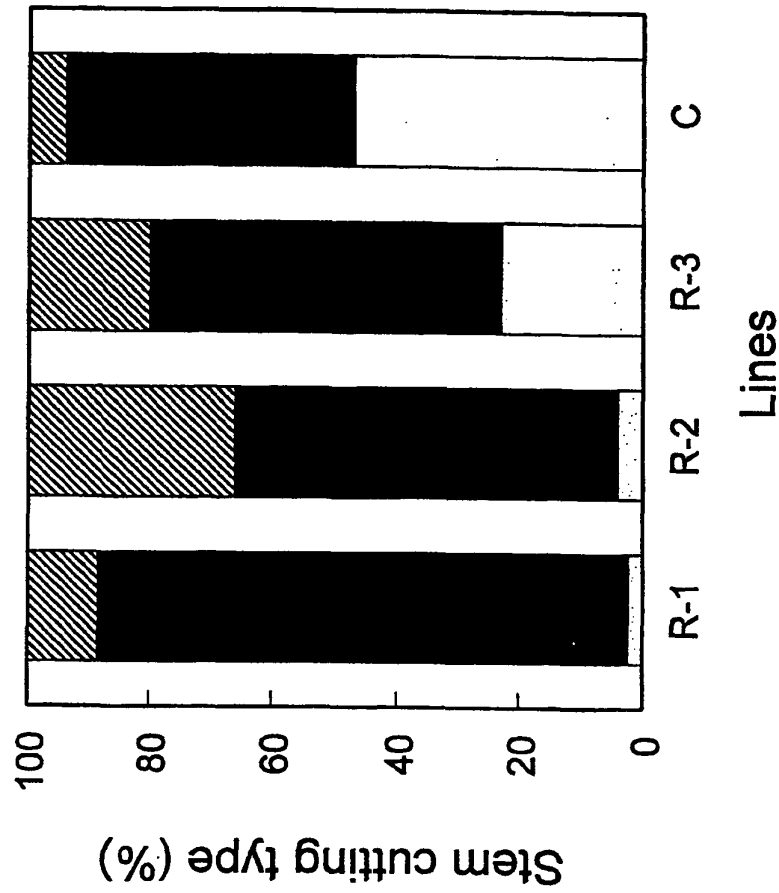


Fig. 9

11/18

206210 "SHHHEO

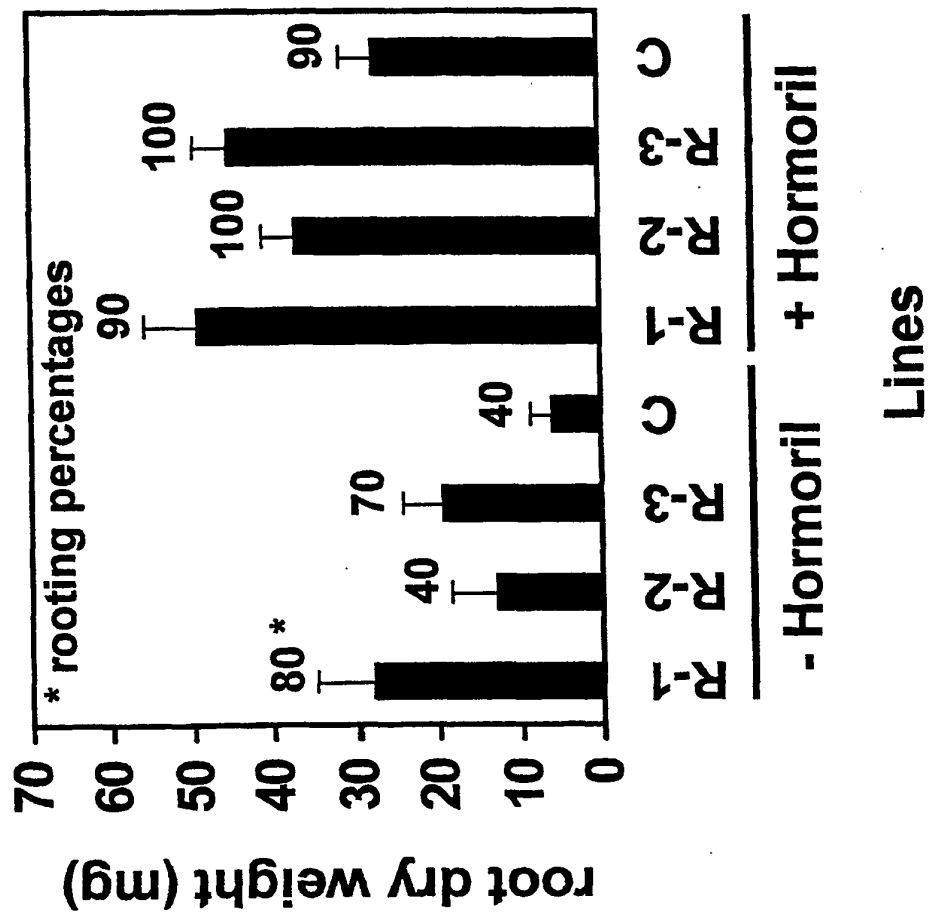


Fig. 10

12/18

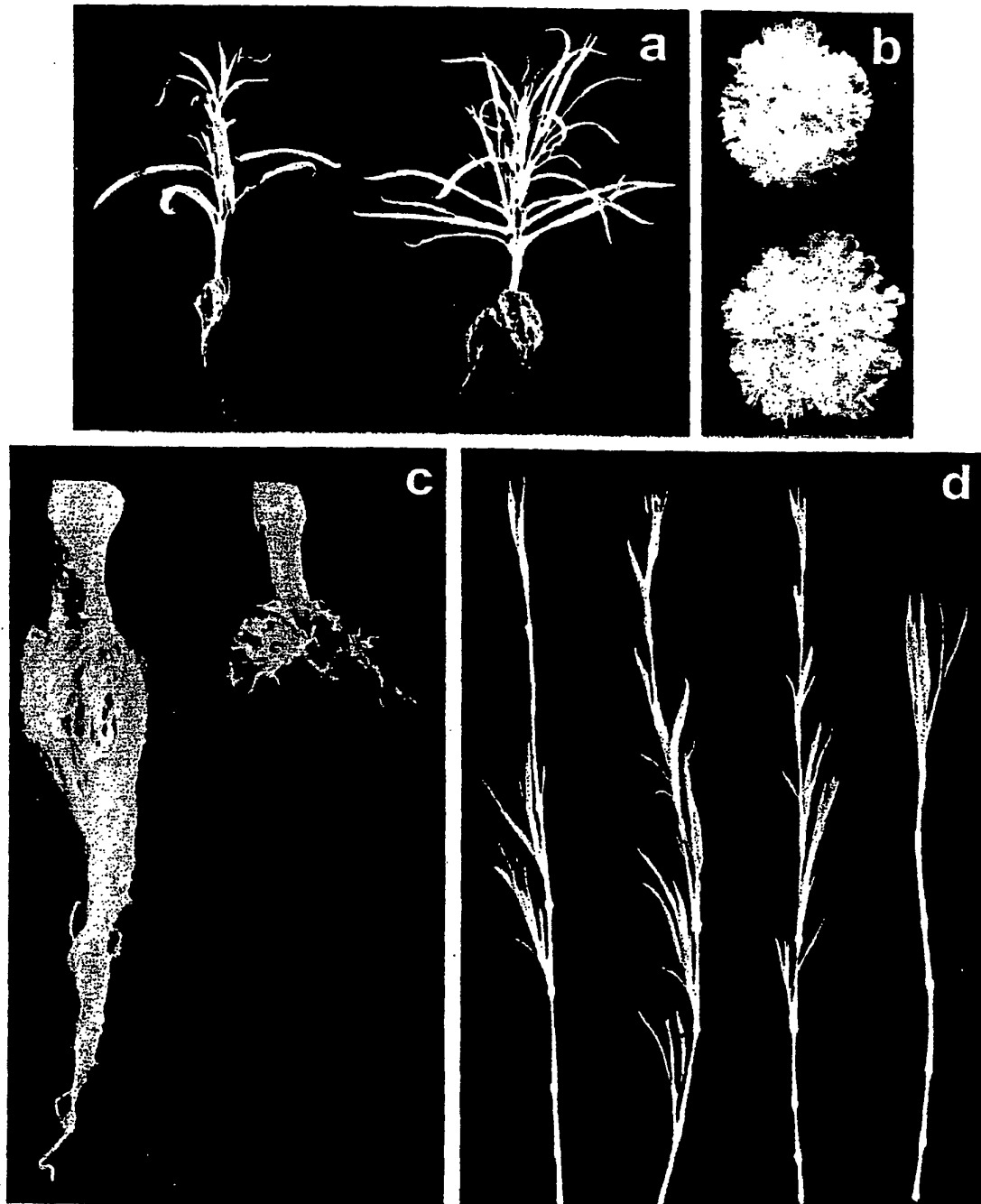


Fig. 11

09/914146-012902

13/18

09914146-012900
206270-94747660

1 ACTATATCTT AAATATTCAC AACATTATAA CATAAGCTTC AAAATAACAT
51 TATTCCGATA TTTACGTAAT ATAATACGTA TCATATTAGG GTACATTCAT
101 TTTATCAACT ACGACTGCAT ATTGTTAGAC AGTCTCATAT ATACGCATAA
151 AAAATGGTCG CTGAAAAACC CAAAACGCTC ACTTCACTAG AAGGGGACGA
201 TAAATTGAAC TCGAATTTTG TTAGGGACGA GGATGAACGT CCGAAAGTGG
251 CGTATAATGA GTTTAGCAAT GATATTCCGG TGATATCTCT TGCTGGTATA
301 GATGGTGAAA AAAGGGGTGA AATATGTCGG AAGATTGTTG AGGCGTGTGA
351 AGATTGGGGG ATTTTCAAG TGGTTGATCA CGGTGTGGT GACGATCTTA
401 TTGCTGATAT GACTCGGTTG GCTCGTGAAT TTTTCGCTCT CCCGGCAGAA
451 GAGAAGCTCC GATTTGATAT GTCTGGTGGT AAAAAGGGCG GTTTTATCGT
501 GTCGAGTCAT CTTCAAGTTCAATCAACAGG GAGAAGTAGT GCAGGACTGG
551 AGGGAAATCGTGACGTATTT CTCATACCCG ACGAACTCAA GGGACTACAC
601 AAGATGGCCAGACAAACCAG AGGGTTGGAT AAAGGTCACA GAGGAATACA
651 GCAACAAGTTAATGACCTTA GCATGTACAC TTTTAGGTGT ACTTTCTGAA
701 GCCATGGGTTTAGAATTAGA GGCACCTACT AAAGCTTGTG TTGATATGGA
751 CCAAAGATTGTGGTTAATT ACTACCCTAA GTGCCCTCAA CCTGACCTTA
801 CTTTAGGGCTCAAGAGGCAC ACCGACCCCG GGACTATAAC CCTCCTCCTT
851 CAGGACCAAGTCGGCGGTCT TCAGGCCACT CGTGACGGTG GTAAACTTG
901 GATTACCGTG CAGCCGGTTC CCGGTGCCTT CGTTGTAAAC CTTGGTGATC
951 ATGGTCATGTTGCGCGAAAA ATGGCCAAAG ACCTTGAGAT CGCCCGTCAT
1001 AAGAGGCTTG CTAAAGAGGA AATGCCTTTT AAAGAGTTGG ACGAGGCCAA
1051 GTTTGAGTCC AAATCTATTG ACCAAATACT TGCTTAGATG GGCTTGGTTT
1101 GGTTTCATTA TATTAAATTT ATTATTATTA TTATTTATTG CATTTGATAT
1151 GATATGATTG GAAATAAAAG AGAGATTGTT TGTGATAATT TGTGTGATTA
1201 TTATATCACT AAGTTATGGC TTTAATTTGT GGTATGTTGG GAATTATATA
1251 TTTAGTTTTG TGTGAAGAAT ATATGATTTA AAGTTAAAAA AAAAAATGAT
1301 TTGTTATATGATTTACTTGT AAGGTTATAA GGTATATTTT ATTGTTGAG
1351 TTTGCGTATA

Fig. 12

206210" 94444650

14/18

1 CCAAGCCCAT CTAAGCAAGT ATTGGTCAA TAGATTGGGA CTCAAACCTTG GCCTCGTCCA
61 ACTCTTTAAA AGGCATTTTC TCTTTAGCAA GCCTCTTATG ACGGCGGATC TCAAGGTCTT
121 TGGCCATTTT TCGCCGAACA TGACCATGAT CACCAAGGTT AACAAACGAAG GCACCGGGAA
181 CCGGCTGCAC GGTAATCCAA GTTTIACCAC CGTCACGAGT GGCCTGAAGA CCGCCGACTT
241 GGTCCTGAAG GAGGAGGGTT ATAGTCCCGG GGTCGGGTGTG CCTCTTGAGC CCTAAAGTAA
301 GGTCAGGTG AGGGCACTTA GGTAGTAAT TAACCACAAT CTTTGGGTCC ATATCAACAC
361 AAGCTTTAGT AAGTGCCCTCT AATTCTAAC CCATGGCTTC AGAAAGTACA CCTAAAAAGTG
421 TACATGCTAA GGTCAATTAA TTGTTGCTGT ATTCTCTGTG GACCTTTATC CAACCCCTCTG
481 GTTTGTCTGG CCATCTTGTG TAGTCCCTTG AGTTCTGTCGG GTATGAGAAA TACGTCAACGA
541 TTTCCCTCCA GTCCTGCACT ACTTCTCCCT GTTGAATTGAA CCTGAAGATG ACTCGACACG
601 ATAAACCCGC CCTTTTACC ACCAGACATA TCAAATCGGA GCTTCTCTTC TGCCGGGAGA
661 GCGAAAAATT CACGAGCCAA CCGAGTCATA TCAGCAATAA GATCGTCACC AACACCGTGA
721 TCAACCACTT GAAAAATCCC CCAATCTTCA CACGCCCTCA CAATCTTCCG ACATATTTCA
781 CCCCTTTTTT CACCATCTAT ACCAGCAAGA GATATCACCG GAATATCAAT GCTAAACTCA
841 TTATACGCCA CTTTCGGACG TTCAATCCTCG TCCCTAACAA AATTCGAGTT CAATTTATCG
901 TCCCCTTCTA GTGAAGTGAG CGTTTGTGG

Fig. 13

15/18

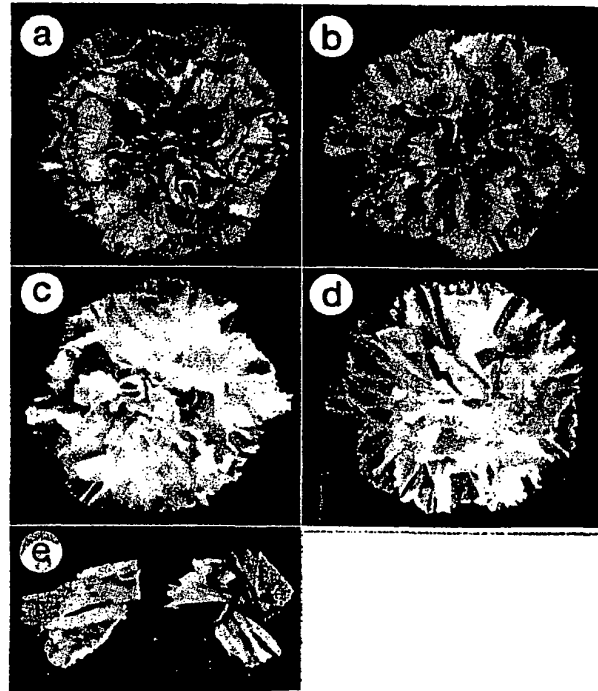


Fig. 14

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16/18

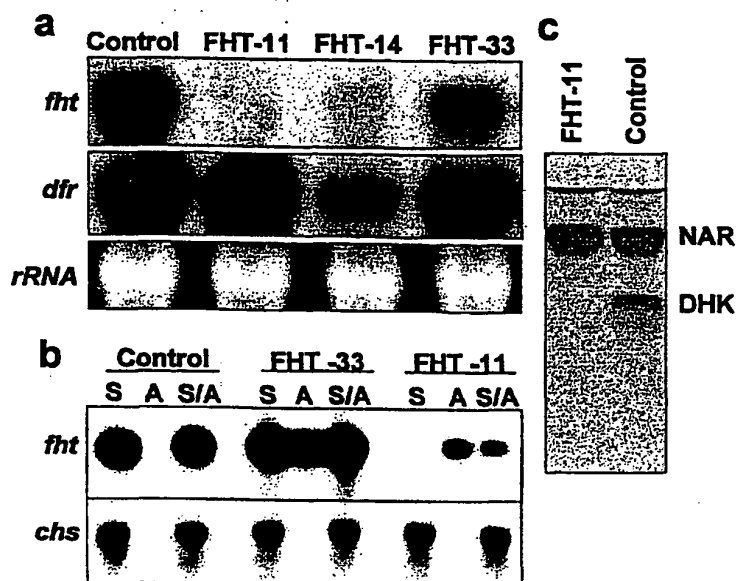


Fig. 15

17/18

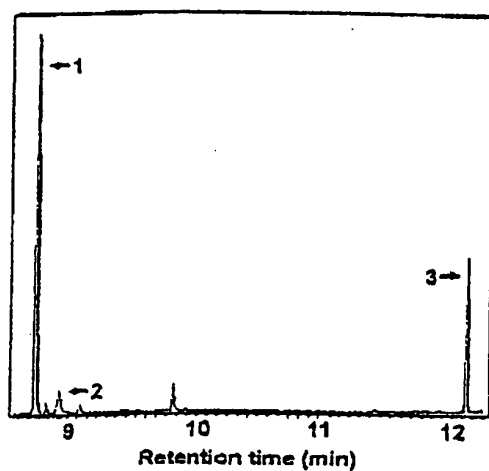


Fig. 16

09/914146-012900

18/18

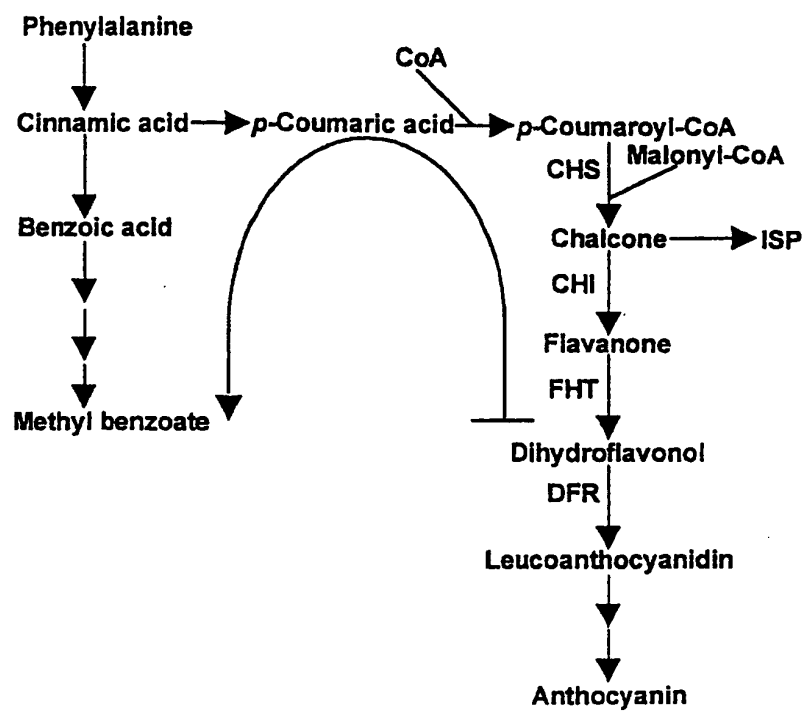


Fig. 17